SEQUENCE LISTING

<110> Reactive Surfaces, Ltd.

<120> Recombinant Organophosphorus Acid Anhydrase and Methods of Use

<130> RACT-00100

<140> US 08/252,384 A
<141> 2004-06-01

<150> 07/928,540
<151> 1992-08-13

<150> 08/252,384
<151> 1994-06-01

<150> 07/344,258
<151> 1989-04-27

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aca atc tct gaa gcg ggt ttc aca ctg act cac gag cac atc tgc ggc       96
Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu His Ile Cys Gly
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agc tcg gca gga ttc ttg cgt gct tgg cca gag ttc ttc ggt agc cgc      144
Ser Ser Ala Gly Phe Leu Arg Ala Trp Pro Glu Phe Phe Gly Ser Arg
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aaa gct cta gcg gaa aag gct gtg aga gga ttg cgc cgc gcc aga gcg      192
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gct ggc gtg cga acg att gtc gat gtg tcg act ttc gat atc ggt cgc      240
Ala Gly Val Arg Thr Ile Val Asp Val Ser Thr Phe Asp Ile Gly Arg
65                  70                  75                  80

gac gtc agt tta ttg gcc gag gtt tcg cgg gct gcc gac gtt cat atc      288
Asp Val Ser Leu Leu Ala Glu Val Ser Arg Ala Ala Asp Val His Ile
                85                  90                  95

gtg gcg gcg acc ggc ttg tgg ttc gac ccg cca ctt tcg atg cga ttg      336
Val Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu
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Arg Ser Val Glu Glu Leu Thr Gln Phe Phe Leu Arg Glu Ile Gln Tyr
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aca ggc aag gcg acc ccc ttt cag gag tta gtg tta aag gcg gcc gcc      480
Thr Gly Lys Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala
145                 150                 155                 160

cgg gcc agc ttg gcc acc ggt gtt ccg gta acc act cac acg gca gca      528
Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala
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agt cag cgc gat ggt gag cag cag gcc gcc att ttt gag tcc gaa ggc      576
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        195                 200                 205

ttg agc tat ctc acc gcc ctc gct gcg cgc gga tac ctc atc ggt cta      672
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    210                 215                 220

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tca gcc ctc ctg ggc atc cgt tcg tgg caa aca cgg gct ctc ttg atc      768
Ser Ala Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile
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aag gcg ctc atc gac caa ggc tac atg aaa caa atc ctc gtt tcg aat      816
Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln Ile Leu Val Ser Asn
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gac tgg ctg ttc ggg ttt tcg agc tat gtc acc aac atc atg gac gtg      864
Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val
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atg gat cgc gtg aac ccc gac ggg atg gcc ttc att cca ctg aga gtg      912
Met Asp Arg Val Asn Pro Asp Gly Met Ala Phe Ile Pro Leu Arg Val
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305                 310                 315                 320

atc act gtg act aac ccg gcg cgg ttc tat gtc acc gac ctt gcg ggc     1008
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Ser Ser Ala Gly Phe Leu Arg Ala Trp Pro Glu Phe Phe Gly Ser Arg
        35                  40                  45

Lys Ala Leu Ala Glu Lys Ala Val Arg Gly Leu Arg Arg Ala Arg Ala
    50                  55                  60

Ala Gly Val Arg Thr Ile Val Asp Val Ser Thr Phe Asp Ile Gly Arg
65                  70                  75                  80

Asp Val Ser Leu Leu Ala Glu Val Ser Arg Ala Ala Asp Val His Ile
                85                  90                  95

Val Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu
            100                 105                 110

Arg Ser Val Glu Glu Leu Thr Gln Phe Phe Leu Arg Glu Ile Gln Tyr
        115                 120                 125

Gly Ile Glu Asp Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr
    130                 135                 140

Thr Gly Lys Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala
145                 150                 155                 160

Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala
                165                 170                 175

Ser Gln Arg Asp Gly Glu Gln Gln Ala Ala Ile Phe Glu Ser Glu Gly
            180                 185                 190

Leu Ser Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp Asp
        195                 200                 205

Leu Ser Tyr Leu Thr Ala Leu Ala Ala Arg Gly Tyr Leu Ile Gly Leu
    210                 215                 220

Asp His Ile Pro His Ser Ala Ile Gly Leu Glu Asp Asn Ala Ser Ala
225                 230                 235                 240

Ser Ala Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile
                245                 250                 255
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Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln Ile Leu Val Ser Asn
            260                 265                 270

Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val
        275                 280                 285

Met Asp Arg Val Asn Pro Asp Gly Met Ala Phe Ile Pro Leu Arg Val
    290                 295                 300

Ile Pro Phe Val Arg Glu Lys Gly Val Pro Gln Glu Thr Leu Ala Gly
305                 310                 315                 320

Ile Thr Val Thr Asn Pro Ala Arg Phe Tyr Val Thr Asp Leu Ala Gly
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Val
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